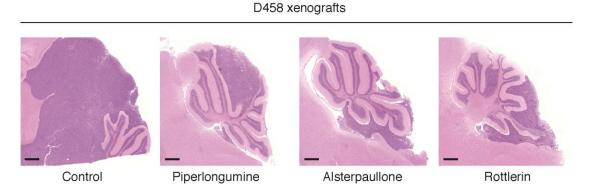
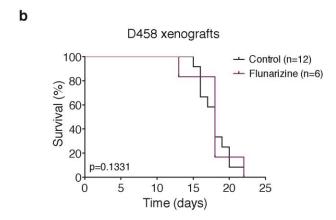
Identification of alsterpaullone as a novel small molecule inhibitor to target group 3 medulloblastoma

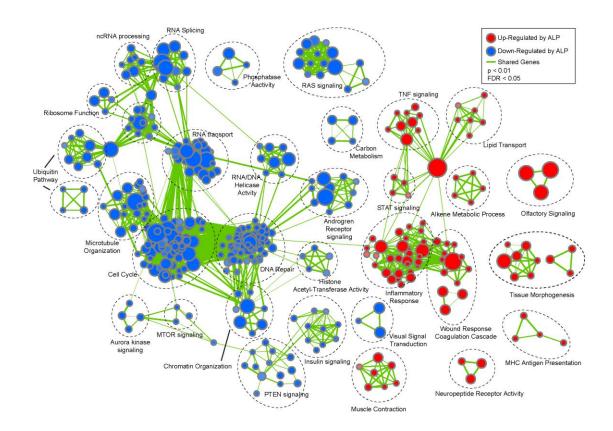
Supplementary Material

а





Supplementary Figure 1. **a** Representative H&E staining demonstrates that mice bearing D458 tumors treated with piperlongumine, alsterpaullone and rottlerin have smaller medulloblastomas in the cerebellum. Scale bar: $500 \, \mu m$. **b** Kaplan-Meier survival curves of D458 medulloblastoma xenografts treated with flunarizine ($50 \, mg/Kg$, daily for 2 weeks; n = 6) or vehicle control (10% DMSO; n = 12). Survival differences were calculated using a log-rank test.

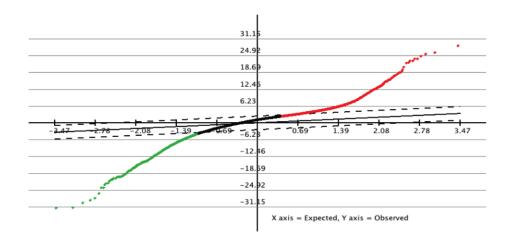


Supplementary Figure 2. Biological pathways and processes up- and down-regulated by alsterpaullone. Gene Set Enrichment Analysis (GSEA) comparing gene sets up- and down-regulated by alsterpaullone (ALP) in D458 and D425 medulloblastoma cells (FDR < 0.05; p < 0.01). Cytoscape and Enrichment Map were used for visualization of the GSEA results. The enriched gene sets were grouped by their similarity, represented as nodes, and mapped as a network. The size of each node determines the total number of genes within each gene set.

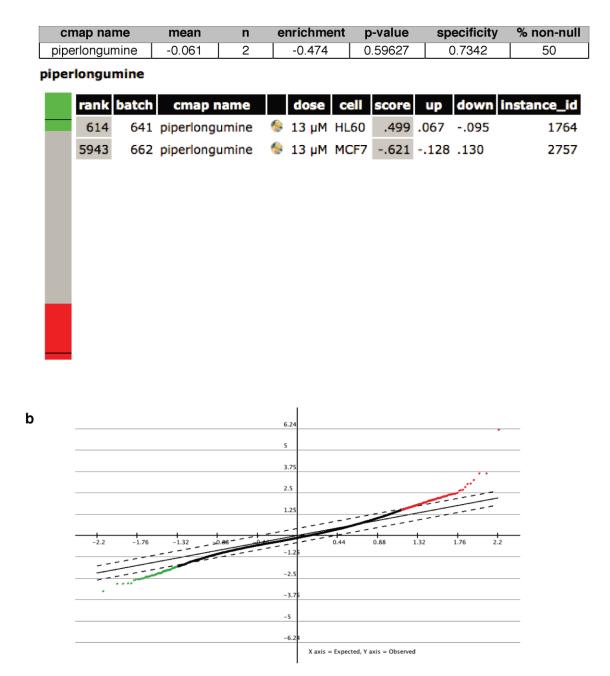
ı	cmap name	mean	n	enrichment	p-value	specificity	% non-null
	alsterpaullone	0.853	3	0.995	< 0.000001	0.0116	100
	alsterpaullone						

ran	k b	atch	cmap name		dose	cell	score	up	down	instance_id
	4	1066	alsterpaullone	-	10 μΜ	MCF7	.934	.248	366	7051
	9	1067	alsterpaullone	8	10 μΜ	PC3	.904	.208	385	7056
3	1	1073	alsterpaullone	8	10 μΜ	PC3	.720	.146	327	7078

b



Supplementary Figure 3. Alsterpaullone reverses Group 3 medulloblastoma gene expression signature as shown by (a) the C-MAP analysis and (b) the Significance Analysis of Microarrays (SAM) plot.



Supplementary Figure 4. The gene expression profile of Group 3 medulloblastomas is not affected by piperlongumine as determined by (a) the C-MAP analysis and (b) the Significance Analysis of Microarrays (SAM) plot.